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AMENDMENTS TO THE CLAIMS

This listing of claims will replace all prior versions, and listings, of claims in the application:

Listing of Claims:

Claims 1-30 (cancelled)

Claim 31 (currently amended): An isolated polynucleotide comprising:

- (a) a nucleotide sequence encoding a lysyl-tRNA synthetase, wherein the amino acid sequence of the synthetase and the amino acid sequence of SEQ ID NO:10 [[have]]share at least 80% identity based on the Clustal alignment method, or
- (b) a full-length complement of the nucleotide sequence of (a).

Claim 32 (currently amended): The polynucleotide of claim 31, wherein the amino acid sequence of the synthetase and the amino acid sequence of SEQ ID NO:10 [[have]]share at least 85% identity based on the Clustal alignment method.

Claim 33 (currently amended): The polynucleotide of claim 31, wherein the amino acid sequence of the synthetase and the amino acid sequence of SEQ ID NO:10 [[have]]share at least 90% identity based on the Clustal alignment method.

Claim 34 (currently amended): The polynucleotide of claim 31, wherein the amino acid sequence of the synthetase and the amino acid sequence of SEQ ID NO:10 [[have]]share at least 95% identity based on the Clustal alignment method.

Claim 35 (previously presented): The polynucleotide of claim 31 comprising the nucleotide sequence of SEQ ID NO:9.

Claim 36 (previously presented): The polynucleotide of claim 31, wherein the synthetase comprises the amino acid sequence of SEQ ID NO:10.

Claim 37 (previously presented): A chimeric gene comprising a polynucleotide of claim 31 operably linked to a regulatory sequence.

Claim 38 (previously presented): A vector comprising a polynucleotide of claim 31.

Claim 39 (previously presented): A method for transforming a cell comprising

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Claim 40 (previously presented): A cell comprising a chimeric gene of claim 37.

Claim 41 (previously presented): A method for producing a plant comprising transforming a plant cell with a chimeric gene of claim 37 and regenerating a plant from the transformed plant cell.

Claim 42 (previously presented): A plant comprising a chimeric gene of claim 37.

Claim 43 (previously presented): A seed comprising a chimeric gene of claim 37.

Claim 44 (withdrawn): An isolated polynucleotide comprising:
(a) a nucleotide sequence encoding an lysyl-tRNA synthetase, wherein the amino acid sequence of the synthetase and the amino acid sequence of SEQ ID NO:16 have at least 90% identity based on the Clustal alignment method, or
(b) the complement of the nucleotide sequence.

Claim 45 (withdrawn): The polynucleotide of claim 44, wherein the amino acid sequence of the synthetase and the amino acid sequence of SEQ ID NO:16 have 95% identity based on the Clustal alignment method.

Claim 46 (withdrawn): The polynucleotide of claim 44 comprising the nucleotide sequence of SEQ ID NO:15.

Claim 47 (withdrawn): The polynucleotide of claim 44, wherein the synthetase comprises the amino acid sequence of SEQ ID NO:16.

Claim 48 (withdrawn): A chimeric gene comprising the polynucleotide of claim 44 operably linked to a regulatory sequence.

Claim 49 (withdrawn): A vector comprising the polynucleotide of claim 44.

Claim 50 (withdrawn): A method for transforming a cell comprising transforming a cell with the polynucleotide of claim 44.

Claim 51 (withdrawn): A cell comprising the chimeric gene of claim 48.

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Claim 52 (withdrawn): A method for producing a plant comprising transforming a plant cell with the chimeric gene of claim 44 and regenerating a plant from the transformed plant cell.

Claim 53 (withdrawn): A plant comprising the chimeric gene of claim 48.

Claim 54 (withdrawn): A seed comprising the chimeric gene of claim 48.

Claim 55 (withdrawn): An isolated polynucleotide comprising:

- (a) a nucleotide sequence encoding an lysyl-tRNA synthetase, wherein the amino acid sequence of the synthetase and the amino acid sequence of SEQ ID NO:12 have at least 85% identity based on the Clustal alignment method, or
- (b) the complement of the nucleotide sequence.

Claim 56 (withdrawn): The polynucleotide of claim 55, wherein the amino acid sequence of the synthetase and the amino acid sequence of SEQ ID NO:12 have 90% identity based on the Clustal alignment method.

Claim 57 (withdrawn): The polynucleotide of claim 55, wherein the amino acid sequence of the synthetase and the amino acid sequence of SEQ ID NO:12 have 95% identity based on the Clustal alignment method.

Claim 58 (withdrawn): The polynucleotide of claim 55 comprising the nucleotide sequence of SEQ ID NO:11.

Claim 59 (withdrawn): The polynucleotide of claim 55, wherein the synthetase comprises the amino acid sequence of SEQ ID NO:12.

Claim 60 (withdrawn): A chimeric gene comprising the polynucleotide of claim 55 operably linked to a regulatory sequence.

Claim 61 (withdrawn): A vector comprising the polynucleotide of claim 55.

Claim 62 (withdrawn): A method for transforming a cell comprising transforming a cell with the polynucleotide of claim 55.

Claim 63 (withdrawn): A cell comprising the chimeric gene of claim 60.

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Claim 64 (withdrawn): A method for producing a plant comprising transforming a plant cell with the chimeric gene of claim 55 and regenerating a plant from the transformed plant cell.

Claim 65 (withdrawn): A plant comprising the chimeric gene of claim 60.

Claim 66 (withdrawn): A seed comprising the chimeric gene of claim 60.